PCT09

RAW SEQUENCE LISTING

DATE: 09/24/2001

PATENT APPLICATION: US/09/807,345

TIME: 16:35:06

Input Set : A:\ES.txt

Output Set: N:\CRF3\09242001\I807345.raw

3 <110> APPLICANT: ARKHAMMAR, Per O.G. et al. 5 <120> TITLE OF INVENTION: AN IMPROVED METHOD FOR EXTRACTING QUANTITATIVE INFORMATION RELATING TO AN

INFLUENCE IN A CELLULAR RESPONSE 8 <130> FILE REFERENCE: 0459-0571P 10 <140> CURRENT APPLICATION NUMBER: 09/807,345 11 <141> CURRENT FILING DATE: 2001-04-12 13 <160> NUMBER OF SEQ ID NOS: 40 15 <170> SOFTWARE: PatentIn version 3.1

ENTERED

17 <210> SEQ ID NO: 1 18 <211> LENGTH: 1788 19 <212> TYPE: DNA

20 <213> ORGANISM: Aequorea victoria and mouse

22 <220> FEATURE:

23 <221> NAME/KEY: CDS

24 <222> LOCATION: (1)..(1788)

25 <223> OTHER INFORMATION: 28 <400> SEQUENCE: 1 29 atg ggc aac gcc gcc gcc aag aag ggc agc gag cag gag agc gtg 48 30 Met Gly Asn Ala Ala Ala Lys Lys Gly Ser Glu Gln Glu Ser Val 33 aaa gag ttc cta gcc aaa gcc aag gaa gat ttc ctg aaa aaa tgg gaa 96 34 Lys Glu Phe Leu Ala Lys Ala Lys Glu Asp Phe Leu Lys Lys Trp Glu 20 25 37 gac ccc tct cag aat aca gcc cag ttg gat cag ttt gat aga atc aag 38 Asp Pro Ser Gln Asn Thr Ala Gln Leu Asp Gln Phe Asp Arg Ile Lys 40 192 41 acc ctt ggc acc ggc tcc ttt ggg cga gtg atg ctg gtg aag cac aag 42 Thr Leu Gly Thr Gly Ser Phe Gly Arg Val Met Leu Val Lys His Lys 55 60 240 45 gag agt ggg aac cac tac gcc atg aag atc tta gac aag cag aag gtg 46 Glu Ser Gly Asn His Tyr Ala Met Lys Ile Leu Asp Lys Gln Lys Val 47 65 70 49 gtg aag cta aag cag atc gag cac act ctg aat gag aag cgc atc ctg 288 50 Val Lys Leu Lys Gln Ile Glu His Thr Leu Asn Glu Lys Arg Ile Leu 53 cag gcc gtc aac ttc ccg ttc ctg gtc aaa ctt gaa ttc tcc ttc aag 336 54 Gln Ala Val Asn Phe Pro Phe Leu Val Lys Leu Glu Phe Ser Phe Lys 105 57 gac aac tca aac ctg tac atg gtc atg gag tat gta gct ggt ggc gag 384 58 Asp Asn Ser Asn Leu Tyr Met Val Met Glu Tyr Val Ala Gly Gly Glu 120 115 432 61 atg ttc tcc cac cta cgg cgg att gga agg ttc agc gag ccc cat gcc 62 Met Phe Ser His Leu Arg Arg Ile Gly Arg Phe Ser Glu Pro His Ala

135

150

65 cgt ttc tac gcg gcg cag atc gtc ctg acc ttt gag tat ctg cac tcc

66 Arg Phe Tyr Ala Ala Gln Ile Val Leu Thr Phe Glu Tyr Leu His Ser

140

155

480

67 145





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							ttg Leu										624
	Glu						ggc Gly 215										672
86							gag Glu										720
							atc Ile										768
							agc Ser		_	_	_	_	_	_			816
							acc Thr										864
	Val		Asp					Lys					Thr			att	912
106	gcc	ato	tat	cag			gtg			ccc	ttc	ata	сса	aad	ttt	aaa	960
107		Ile		Gln	Arg	Lys 310		Glu	Ala	Pro	Phe 315	Ile				Lys 320	
109	Ala 305 ggc Gly	Ile	Tyr ggg	gac	acg	310 agt Ser	aac	ttt	. gac	gac	315 tat Tyr	: Ile . gag	Pro	Lys gaa	Phe gag	Lys 320 atc Ile	1008
109 110 111 113	Ala 305 ggc Gly cgg Arg	cct Pro	ggg Gly	gac Asp	acg Thr 325 aat	310 agt Ser	aac Asn aag	ttt Phe tgt	gac Asp	gac Asp 330 aag	315 tat Tyr gag	gag Glu	gag Glu act	gaa Glu gag	gag Glu 335 ttt	Lys 320 atc Ile	
109 110 111 113 114 115 117	Ala 305 ggc Gly cgg Arg	cct Pro gtc Val	ggg Gly tcc Ser	gac Asp atc Ile 340 agt	acg Thr 325 aat Asn	310 agt Ser gag Glu	aac Asn aag Lys	ttt Phe tgt Cys	gac Asp Gly 345 ctt	gac Asp 330 aag Lys	315 tat Tyr gag Glu	gag Glu ttt Phe	gag Glu act Thr	gaa Glu gag Glu 350 gtc	gag Glu 335 ttt Phe	Lys 320 atc Ile ggg	•
109 110 111 113 114 115 117 118 119 121	Ala 305 ggc Gly cgg Arg cgc Arg	cct Pro gtc Val gcc Ala	ggg Gly tcc Ser atg Met 355 gaa Glu	gac Asp atc Ile 340 agt Ser	acg Thr 325 aat Asn aaa Lys	310 agt Ser gag Glu gga Gly	aac Asn aag Lys gaa Glu	ttt Phe tgt Cys gaa Glu 360 gtt	gac Asp ggc Gly 345 ctt Leu	gac Asp 330 aag Lys ttc Phe	315 tat Tyr gag Glu act Thr	gag Glu ttt Phe gga Gly	gag Glu act Thr gtt Val 365 ttc	gaa Glu gag Glu 350 gtc Val	gag Glu 335 ttt Phe cca Pro	Lys 320 atc Ile ggg Gly att Ile	1056
109 110 111 113 114 115 117 118 119 121 122 123 125 126	Ala 305 ggc Gly cgg Arg ctt Leu gga Gly	gtc Val gcc Ala gtt Val 370 gag Glu	ggg Gly tcc Ser atg Met 355 gaa Glu	gac Asp atc Ile 340 agt Ser tta Leu	acg Thr 325 aat Asn aaa Lys gat Asp	310 agt Ser gag Glu gga Gly ggc Gly	aac Asn aag Lys gaa Glu gat Asp 375 gca Ala	ttt Phe tgt Cys gaa Glu 360 gtt Val	gac Asp ggc Gly 345 ctt Leu aat Asn	gac Asp 330 aag Lys ttc Phe Gly	315 tat Tyr gag Glu act Thr	gag Glu ttt Phe gga Gly aaa Lys 380 ctt	gag gag Glu act Thr yal 365 ttc Phe	gaa Glu gag Glu 350 gtc Val tct	gag Glu 335 ttt Phe cca Pro	Lys 320 atc Ile ggg Gly att Ile agt	1056 1104
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135	222	020	aa+		+++	++0	aag	ant		a+a	000	422	aat		ata	cac	1344
		_		-			Lys	-	-	-		-			_	-	Tlaa
139	пÃ2	GIII	435	мэр	FIIE	File	пуз	440	мта	Het	PIO	Giu	445	ıyı	Val	GIII	
	~~~	242		2+2	+++	+20	aaa		<b>424</b>	aaa	220	tag		202	aat	act	1392
	-	_					Lys	_	_				_		-	-	1392
143	GIU	450	TIIT	TIE	Pile	тут	цуS 455	ASP	ASP	СТУ	ASII	460	гуз	1111	AIY	AIG	
	as s		220	+++	maa.	aat	gat	200	ott	att	aat		ato	man	tta	222	1440
							Asp										1440
	465	Val	цуз	FIIE	GIU	470	кар	1111	пец	Val	475	лту	116	GIU	пеп	480	
		2++	a a t	+++	222		gat	aaa	aac	att		mma	cac	222	ata		1488
			_			-	Asp								-	-	1400
151	СТУ	116	кор	FIIC	485	GIU	кор	GLY	LOII	490	пец	GIY	птэ	цуз	495	Giu	
	tac	22+	tat	aac		cat	aat	αta	tac		ato	aca	gac	222		aan	1536
							Asn										1330
155	1 Y 1	Maii	1 Y 1	500	Der	1113	ASII	Vai	505	110	IIC C	AIU	пор	510	110	цуб	
	aat	σσο	atc		att	aac	ttc	aaa		ana	cac	aac	att		nat	σσα	1584
					-		Phe			_					_		1301
159	UPII	GLY	515	цуз	Vai	ASII	riic	520	110	nry	1113	AJII	525	цуз	иор	GIY	
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163	DCI	530	02.11	нси	1114	1155	535	-1-	0111	01	11011	540			011		
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	Lys	Glu	Phe	Leu	Ala	Lys	Ala	Lys	Glu	Asp	Phe	Leu	Lys	Lys	$\mathtt{Trp}$	Glu	
194				20					25					30			
197	Asp	Pro	Ser	Gln	Asn	Thr	Ala	Gln	Leu	Asp	Gln	Phe	Asp	Arg	Ile	Lys	
198			35					40					45				
201	Thr	Leu	Gly	Thr	Gly	Ser	Phe	Gly	Arg	Val	Met	Leu	Val	Lys	His	Lys	
202		50					55					60					
205	Glu	Ser	Gly	Asn	His	$\mathtt{Tyr}$	Ala	Met	Lys	Ile	Leu	Asp	Lys	Gln	Lys	Val	



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	Gln	Ala	Val	Asn	Phe	Pro	Phe	Leu	Val	Lys	Leu	Glu	Phe	Ser	Phe	Lys
214				100					105	-				110		-
	Asp	Asn	Ser	Asn	Leu	Tyr	Met	Val	Met	Glu	Tyr	Val	Ala	Gly	Gly	Glu
218	-		115			•		120			-		125	_	-	
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222		130				_	135		-	_		140				
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	145		-			150					155		-			160
229	Leu	Asp	Leu	Ile	Tyr	Arq	Asp	Leu	Lys	Pro	Glu	Asn	Leu	Leu	Ile	Asp
230		•			165		-	•	-	170					175	-
233	Gln	Gln	Gly	Tyr	Ile	Gln	Val	Thr	Asp	Phe	Gly	Phe	Ala	Lys	Arq	Val
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242		210				-	215	-		-		220	-	-	-	
245	Leu	Gly	Val	Leu	Ile	Tyr	Glu	Met	Ala	Ala	Gly	Tyr	Pro	Pro	Phe	Phe
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273	Arg	Val	Ser	Ile	Asn	Glu	Lys	Cys	_	Lys	Glu	Phe	Thr	Glu	Phe	Gly
274				340					345					350		
	Arg	Ala	Met	Ser	Lys	Gly	Glu	Glu	Leu	Phe	Thr	Gly	Val	Val	Pro	Ile
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310					485					490					495		
	Tyr	Asn	Tyr		Ser	His	Asn	Val	_	Ile	Met	Ala	Asp	_	Pro	Lys	
314	λan	Cl v	Tla	500	Va 1	λan	Dha	T.ve	505	Δra	Uic	λen	Ile	510	λen	Gl _W	
318	ASII	Сту	515	цуз	Val	Maii	FIIC	520	116	ALY	1113	N3II	525	цуз	кэр	GLY	
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330	nha	37-3	mbs	71.	565	C1	т1а	mh∞	111.0	570	Wot	) an	C1.,	Т ом	575	Tria	
334	Pne	vaı	TIII	580	Ата	СТУ	ire	TILL	585	СТЙ	Met	ASP	Glu	590	TAT	гух	
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/807,345



DATE: 09/24/2001 TIME: 16:35:07

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Output Set: N:\CRF3\09242001\I807345.raw